<u>RAW SEQUENCE LISTING</u>

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _____ 09/123,253A_____

Source: _____ IFW/6, _____

Date Processed by STIC: _____ 05/8/2006_____

ENTERED

CRF Errors Edited by the STIC Systems Branch

Serial Number: _09/123, 253A_   CRF Edit Date: _05/8/2006_
                                 Edited by: _DA_

____ Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line

____ Corrected the SEQ ID NO. Sequence numbers edited were:

_____

____ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

_____

____ Deleted: ___ invalid beginning/end-of-file text ; ___ page numbers

____ Inserted mandatory headings/numeric identifiers, specifically:

_____

____ Moved responses to same line as heading/numeric identifier, specifically:

_____

_✓_ Other: _Insert <220> Mandatory Numeric Identifier_

Revised 09/09/2003

IFW16

RAW SEQUENCE LISTING   DATE: 05/08/2006
PATENT APPLICATION: **US/09/123,253A**   TIME: 10:51:18

Input Set : **A:\pto.da.txt**
Output Set: **N:\CRF4\05052006\I123253A.raw**

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 4 <110> APPLICANT: Yip, Tai-Tung
 5        Hutchens, T. William
 7 <120> TITLE OF INVENTION: Method and Apparatus for Desorption and Ionization of
 8        Analytes
10 <130> FILE REFERENCE: Hutchens D-5639-C4
12 <140> CURRENT APPLICATION NUMBER: 09/123,253A
14 <141> CURRENT FILING DATE: 1998-07-27
16 <150> PRIOR APPLICATION NUMBER: 08/068,896
18 <151> PRIOR FILING DATE: 1993-05-28
20 <160> NUMBER OF SEQ ID NOS: 1
22 <170> SOFTWARE: PatentIn Ver. 2.0
24 <210> SEQ ID NO: 1
26 <211> LENGTH: 26
28 <212> TYPE: PRT
30 <213> ORGANISM: Unknown
32 <220> FEATURE:
35 <223> OTHER INFORMATION: Description of Unknown Organism: peptide
37 <400> SEQUENCE: 1
39 Gly His His Pro His Gly His His Pro His Gly His His Pro His Gly
42   1               5                   10                  15
45 His His Pro His Gly His His Pro His Gly
48              20                  25
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 05/08/2006
PATENT APPLICATION: US/09/123,253A   TIME: 10:51:19

Input Set : **A:\pto.da.txt**
Output Set: **N:\CRF4\05052006\I123253A.raw**

<u>**Invalid Line Length:**</u>
The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 10

VERIFICATION SUMMARY   DATE: 05/08/2006
PATENT APPLICATION: **US/09/123,253A**   TIME: 10:51:19

Input Set : **A:\pto.da.txt**
Output Set: **N:\CRF4\05052006\I123253A.raw**

Raw Sequence Listing before editing,
for reference only

IFW16

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 4 <110> APPLICANT: Yip, Tai-Tung
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Does Not Comply
Corrected Diskette Needed
(pg -1)

ERRORED SEQUENCES

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24 <210> SEQ ID NO: 1
26 <211> LENGTH: 26
28 <212> TYPE: PRT
30 <213> ORGANISM: Unknown
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> Inserted <220>

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W--> 33 .
33 <223> OTHER INFORMATION: Description of Unknown Organism: peptide
35 <400> SEQUENCE: 1
37 Gly His His Pro His Gly His His Pro His Gly His His Pro His Gly
40  1               5                  10                  15
43 His His Pro His Gly His His Pro His Gly
46             20                  25
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 05/08/2006
PATENT APPLICATION: **US/09/123,253A**   TIME: 10:47:51

Input Set : **A:\PATSEQ.txt**
Output Set: **N:\CRF4\05052006\I123253A.raw**

<u>**Invalid Line Length:**</u>
The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 10

VERIFICATION SUMMARY   DATE: 05/08/2006
PATENT APPLICATION: **US/09/123,253A**   TIME: 10:47:51

Input Set : **A:\PATSEQ.txt**
Output Set: **N:\CRF4\05052006\I123253A.raw**

L:33 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:35 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:1